```
                        GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on:        May  4, 2005, 12:06:42 ; Search time 259.813
Seconds
                                    (without alignments)
                                    5365.716 Million cell
updates/sec

Title:         US-09-300-482-27
Perfect score: 229
Sequence:      1
cagaccttattttttctgtc..........cttcaatttatggggaacaa 229


Scoring table:  IDENTITY_NUC
                Gapop 10.0 , Gapext 1.0

Searched:       5642217 seqs, 3043843248 residues

Total number of hits satisfying chosen parameters:      11284434

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries


Database :        Published_Applications_NA:*
                1:  /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
                2:  /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
                3:  /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
                4:  /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
                5:  /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
                6:  /cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
                7:  /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
                8:  /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
                9:  /cgn2_6/ptodata/1/pubpna/US09A_PUBCOMB.seq:*
                10: /cgn2_6/ptodata/1/pubpna/US09B_PUBCOMB.seq:*
                11: /cgn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
                12: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
                13: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
```